

The Chlamydia-Secreted Protease CPAF Promotes Chlamydial Survival in the Mouse Lower Genital Tract

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Despite the extensive *in vitro* characterization of CPAF (chlamydial protease/proteasome-like activity factor), its role in chlamydial infection and pathogenesis remains unclear. We now report that a *Chlamydia trachomatis* strain deficient in expression of CPAF (L2-17) is no longer able to establish a successful infection in the mouse lower genital tract following an intravaginal inoculation. The L2-17 organisms were cleared from the mouse lower genital tract within a few days, while a CPAF-sufficient *C. trachomatis* strain (L2-5) survived in the lower genital tract for more than 3 weeks. However, both the L2-17 and L2-5 organisms maintained robust infection courses that lasted up to 4 weeks when they were directly delivered into the mouse upper genital tract. The CPAF-dependent chlamydial survival in the lower genital tract was confirmed in multiple strains of mice. Thus, we have demonstrated a critical role of CPAF in promoting *C. trachomatis* survival in the mouse lower genital tracts. It will be interesting to further investigate the mechanisms of the CPAF-dependent chlamydial pathogenicity.

Chlamydia trachomatis is a leading cause of sexually transmitted bacterial infection. Following the initial infection in the lower genital tract, the chlamydial organisms can ascend to the upper genital tracts. The upper genital tract infection may cause pathologies such as hydrosalpinx, leading to severe sequelae, including ectopic pregnancy and tubal factor infertility (1, 2). It remains unknown how *C. trachomatis* establishes a successful initial infection in the lower genital tract and achieves ascending infection. The mouse genital tract infection model has been used with either *C. trachomatis* (3–7) or *C. muridarum* (8–15) for studying chlamydial pathogenesis and immunity. *C. trachomatis* infection in the mouse genital tract, unlike *C. muridarum* infection, often fails to induce a long-lasting upper genital tract pathology such as hydrosalpinx. Furthermore, innate immunity alone appeared to be sufficient for controlling the infection (16). Nevertheless, following an intravaginal inoculation, the *C. trachomatis* organisms were found to survive in the lower genital tract for weeks and invasion of the uterine endometrial epithelia was also detected (3, 5). When *C. trachomatis* was directly introduced into the mouse endometrial epithelia via a transcervical or intrauterine inoculation (bypassing the cervical barrier), the organisms were able to induce more-robust inflammatory responses and immunity in the genital tract (7, 17). Thus, mouse genital tract infection with *C. trachomatis* via either intravaginal or intrauterine inoculation has been used for investigating *C. trachomatis* pathogenesis.

Using the *C. trachomatis* genital tract infection mouse model, Sturdevant et al. (3) correlated mutations in the hypothetical open reading frame (ORF) CT135 with the survival/infectivity of *C. trachomatis* serovar D in the mouse genital tracts. However, the mechanisms by which CT135 contributes to the pathogenesis of *C. trachomatis* serovar D remain unknown. Ramsey et al. (6) also used the mouse model for defining the role of the plasmid-encoded Pgp3 in the survival of *C. trachomatis* in the mouse genital tract. This finding was validated in a *C. muridarum* infection mouse model in which a Pgp3-deficient *C. muridarum* strain was no longer able to induce hydrosalpinx (18). Pgp3 is an immuno-

dominant antigen that is secreted into the cytosol of the infected cells (19–21). Many other *C. trachomatis* proteins, including CT311 (22, 23), CT621/622 (24, 25), CT795 (26), and cHtrA (27) as well as GlgA (28), all encoded by hypothetical ORFs, have also been localized in the cytosol of the infected cells. However, the functions of these proteins are largely unknown.

Another well-characterized chlamydial secretion protein is the serine protease CPAF (chlamydial protease/proteasome-like activity factor). Although CPAF was initially discovered as a consequence of its ability to robustly degrade intracellular proteins of the infected cells (29–31), the extent to which these host targets are degraded during infection is a subject of debate (32–37). CPAF is a unique serine protease (38–42) that is secreted into the host cell cytosol via a sec-dependent type II secretion pathway (34, 43). We have recently validated the secretion of CPAF into the host cell cytosol (44). To further investigate the role of CPAF in chlamydial pathogenesis, we took advantage of chemically derived *C. trachomatis* L2 mutants (34, 45, 46) and compared a CPAF-deficient *C. trachomatis* strain to a control strain derived by lateral gene transfer (34) for survival in the mouse genital tract. We found that the CPAF-deficient *C. trachomatis* strain lost its ability to establish a successful infection in the mouse lower genital tract following an intravaginal inoculation whereas its ability to infect the upper genital tract was not significantly affected. This result indicates that

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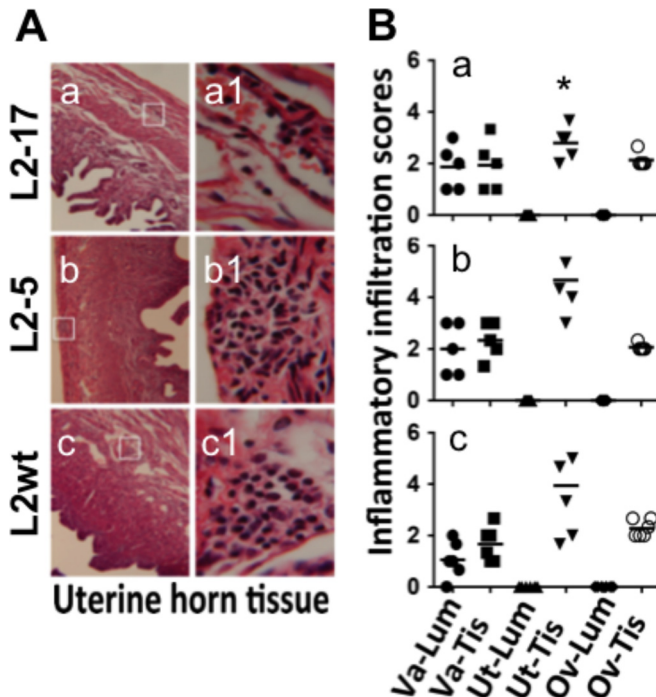
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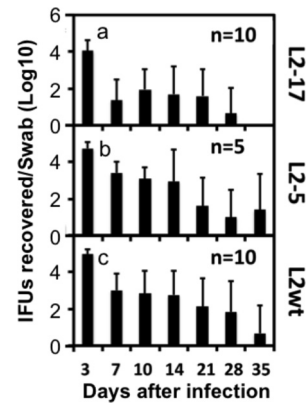
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mice are known to be more resistant to chlamydial infection than C3H/HeJ mice (51). C57BL/6J mice tend to develop Th1-dominant adaptive immunity upon chlamydial infection, while BALB/c mice develop Th2-dominant adaptive immunity (52). We then compared the survival rates seen with L2-17 and L2-5 in the lower genital tracts of C57BL/6J, C3H/HeJ, and BALB/c mice (Fig. 5). The L2-17 organisms failed to survive in the lower genital tracts regardless of the mouse strains tested. The organisms were cleared from the mouse lower genital tracts within a few days of the inoculation. These observations suggest that the lower genital tract factors responsible for clearing L2-17 infection are shared among different mouse strains. To further validate whether the failure of L2-17 to survive in the mouse lower genital tract is mainly due to lack of CPAF, we complemented L2-17 with a plasmid-encoded wild-type CPAF to produce the transformant L2-17/CPAF. We found that the L2-17/CPAF organisms showed significantly in-



creased survival in the mouse lower genital tract (Fig. 6). As a control, the L2-17 organisms were similarly transformed with a mCherry-expressing plasmid. The L2-17/mCherry organisms remained defective in survival in the mouse lower genital tract.

DISCUSSION

In the current report, we have presented the first experimental evidence demonstrating that CPAF promotes chlamydial survival in the mouse lower genital tract but not in the upper. First, the CPAF-deficient *C. trachomatis* L2-17 strain failed to infect the mouse lower genital tract whereas the CPAF-sufficient L2-5 control strain was able to cause a 2-week-long infection, indicating that CPAF plays a critical role in promoting *C. trachomatis* survival in the mouse lower genital tract tissues. Second, the uterine tissue of mice intravaginally inoculated with L2-17 lacked significant inflammatory infiltration whereas the uterine tissue of mice similarly inoculated with L2-5 developed significant inflammatory infiltration, suggesting that L2-17 either failed to reach to the uterine tissue or was unable to provoke uterine inflammation. Third, both L2-17 and L2-5 developed robust infections in the mouse genital tracts and induced significant inflammatory infiltration in the uterine tissue when the organisms were directly delivered into the upper genital tract. These observations have confirmed that CPAF is not essential for the *C. trachomatis* survival and induction of inflammation in the upper genital tracts. The result described above further suggests that lack of uterine inflammation in mice intravaginally infected with L2-17 is probably due to the failure of L2-17 to reach the uterine tissue, since L2-17 is able to activate uterine inflammation when directly delivered to the uterine tissue. Fourth, the requirement for CPAF to aid in chlamydial survival in the mouse lower genital tract was reproduced in 3 different strains of mice despite their different local cytokine profiles and varied adaptive immunity phenotypes (52, 53). Finally, complementation of the L2-17 organisms with a CPAF-expressing plasmid partially restored their

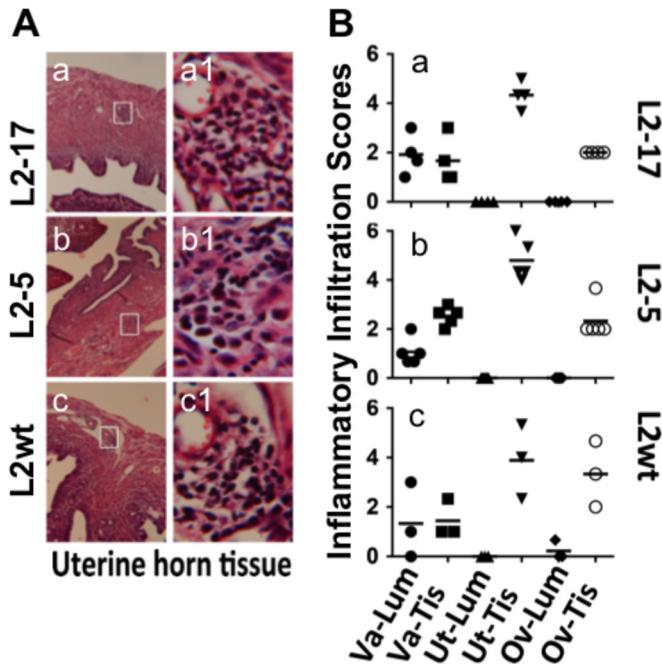


FIG 4 Effect of CPAF deficiency on inflammatory infiltration in the mouse genital tracts following an intrauterine inoculation. (A) The mice described in the Fig. 3 legend were sacrificed on days 21 to 60 postinfection, and the genital tracts were processed for microscopic detection of inflammatory infiltration. Representative images of uterine horn tissue from each group of mice were taken under 10 \times (a to c) and 100 \times (a1 to c1) objective lenses. The areas covered under the 100 \times lens are marked with white rectangles in the corresponding 10 \times images. (B) Different sections of the genital tract, including the lumen (Lum) and tissue (Tis) of vagina (Va), uterine/uterine horns (Ut), and oviducts (Ov), were semiquantitatively scored for inflammatory infiltration based on the criteria described in Materials and Methods. The inflammatory scores from each mouse were plotted individually along the y axis. Solid circles stand for the inflammatory scores from Va-Lum, solid squares for scores from Va-Tis, solid upright triangles for scores from Ut-Lum, solid upside triangles for scores from Ut-Tis, solid diamonds for scores from Ov-Lum, and open circles for scores from Ov-Tis. No significant differences in inflammatory infiltration scores were found between the groups (Wilcoxon).

survival in the mouse lower genital tract whereas a similar transformation performed with a mCherry-expressing control plasmid failed to do so. The significant but partial rescue of the chlamydial infectivity by the CPAF complementation may have been a consequence of either the interference from the excessive amount of the incompletely processed CPAF fragments in the complemented strain (38, 40–42, 44) or a CPAF-independent defect inherited in the parental L2-17 strain (34). Nevertheless, the complementation experiments have demonstrated that the failure of L2-17 to survive in the mouse lower genital tract was at least partially due to the lack of CPAF.

The previously published controversial findings on CPAF degradation of cellular proteins in cell culture systems (32, 33) motivated us to use the L2-17 strain (34) for evaluating the potential functions of CPAF in the mouse genital tracts, which led us to discover an essential role of CPAF in promoting chlamydial survival in the mouse lower genital tract. This mouse model may also allow us to further investigate the mechanisms by which CPAF aids in chlamydial survival. For example, questions on whether CPAF promotes chlamydial ascension from the lower to the upper genital tract and whether CPAF aids in chlamydial infectivity in

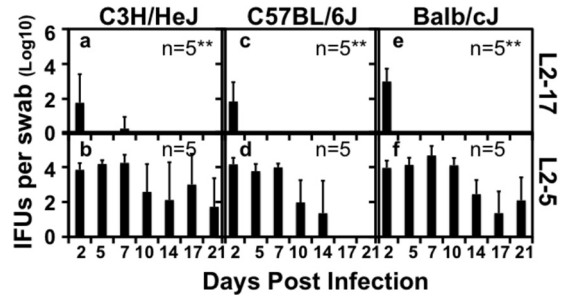


FIG 5 CPAF-dependent chlamydial survival in the lower genital tracts of multiple strains of mice. Six-week-old female C3H/HeJ (a and b; $n = 5$ for each group), C57BL/6J (c and d; $n = 5$), and BALB/cJ (e and f; $n = 5$) mice were intravaginally inoculated with 10^6 IFUs of L2-17 (a, c, and e) or L2-5 (b, d, and f). Vaginal/cervical swabs were taken on different days after the inoculation as indicated along the x axis for monitoring live-organism recovery as shown along the y axis. Note that the L2-17 organisms displayed significantly lower levels of live-organism recovery than the L2-5 organisms regardless of the strains of mice infected (Wilcoxon; **, $P < 0.01$).

different genital tract tissues can now be addressed by simultaneously monitoring the numbers of infectious chlamydial organisms and the numbers of genome copies in multiple genital tract tissues. The next issue is that of how CPAF promotes chlamydial survival/ascension/infectivity. It has been hypothesized that CPAF accumulated in the infected host cell cytosol at the late stage of intracellular chlamydial growth (29) may be released to confront the extracellular mucosal effectors before the intrainclusion organisms are exposed to extracellular environments during host cell lysis and chlamydial spreading (33). The fact that L2-17 was cleared within a few days after inoculation suggests that the innate immunity effectors of the lower genital tract are effective for controlling the infection and that these same effectors may be targeted by CPAF. This hypothesis is consistent with the finding that innate immunity is sufficient for controlling *C. trachomatis* infection in the mouse genital tract (16). The CPAF-dependent chlamydial survival in the mouse lower genital tract was reproduced in 3 different strains of mice, suggesting that CPAF may target host

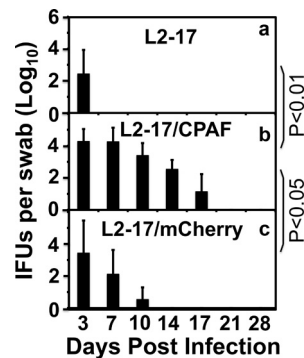


FIG 6 CPAF complementation effect on chlamydial survival in mouse lower genital tract. Seven-week-old C57BL/6J female mice were each inoculated with 1×10^6 IFUs of L2-17 (a; $n = 5$), L2-17/CPAF (b; $n = 5$), or L2-17/mCherry (c; $n = 5$) intravaginally. On different days after inoculation as indicated along the x axis, vaginal swabs were taken for monitoring recovery of live chlamydial organisms. The recovered live organisms were expressed as log₁₀ inclusion-forming units (IFUs) per swab as shown along the y axis. Note that L2-17/CPAF developed more-extensive live-organism shedding than either L2-17 or L2-17/mCherry ($P < 0.01$ or $P < 0.05$, respectively; Wilcoxon).

factors in the lower genital tract shared by the different strains. It is unlikely that the different H-2 haplotypes (C57BL/6), H-2^b, C3H/HeJ, H-2^k, BALB/c, H-2^d) and the other strain-specific factors (51–53) are targeted by CPAF. We recently reported that CPAF selectively degraded cathelicidin LL-37 and neutralized its anti-chlamydial activity (54). It will be interesting to use mice deficient in CRAMP (55), a cathelin-related antimicrobial peptide and the mouse homolog of human LL-37 that is produced in all mouse genital tracts (56), to test whether the CRAMP deficiency can rescue L2-17 in the mouse lower genital tract.

When CPAF was initially discovered, attention was mainly focused on its potential role in dealing with the intracellular proteins of the infected cells (57, 58). Recent studies suggest that CPAF and other chlamydial proteins secreted into the cytoplasm of the infected cells may target extracellular molecules (33, 54). However, caution should be taken in interpreting the interactions of chlamydial factors with host factors detected in *in vitro* systems. The biological relevance of the *in vitro* observations has to be validated using mutant chlamydial organisms in animal models as demonstrated in the current study. As generation of chlamydial mutants becomes easier and animal models continue to be optimized, it is expected that more biologically relevant chlamydial pathogenic mechanisms will be uncovered. A caveat is that some of these chlamydial mutants may be highly attenuated in their growth properties, including attachment, entry, intracellular replication, and exit. The reduced survival/infectivity of the mutants in the mouse genital tract may reflect only their slow/inefficient or defective growth abilities. Indeed, L2-17 produced 3-fold-fewer infectious progeny elementary bodies (EBs) in cultured cells than L2-5 (34), which may have contributed to the significantly shortened survival of L2-17 in the mouse lower genital tract. However, when L2-17 and L2-5 were directly delivered to the upper genital tract, both maintained robust infection courses. Although the infection course of L2-17 was less robust than that of L2-5, probably due to the reduced ability of L2-17 to produce EBs (34), the difference was not significant. Thus, CPAF deficiency preferentially reduced chlamydial survival in the mouse lower but not upper genital tract. We hypothesize that in addition to the growth defect, the L2-17 organisms may also be more susceptible to the lower genital tract effectors. A test of this hypothesis is under way.

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